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6) IMC 129	6) IMC 129	6) IMC 129	6) IMC 129
5) Q508	5) Q508	5) Q508	5) Q508
8) Q4275	8) Q4275	8) Q4275	8) Q4275
Fad2-D wt	Fad2-D wt	Fad2-D wt	Fad2-D wt
Fad2-D (GA316)	Fad2-D (GA316)	Fad2-D (GA316)	Fad2-D (GA316)
Fad2-F wt	Fad2-F wt	Fad2-F wt	Fad2-F wt
Fad2-F (TA515)	Fad2-F (TA515)	Fad2-F (TA515)	Fad2-F (TA515)
Fad2-F (GA908)	Fad2-F (GA908)	Fad2-F (GA908)	Fad2-F (GA908)
ATGGGTGCAGGTGGAATGCAAGTGTCTCCTCCTCCAAATGCAAGTGTCTCCTCCTCCAAATGCAAGTGTCTCCTCCTCCAAATGCAAGTGTCTCCTCCTCCAAATGCAAGTGTCTCCTCCTCCAAATGCAAGTGTCTCCTCCTCCAAATGCAAGTGTCTCCTCCTCCAAATGCAAGTGTCTCCTCCTCCAAATGCAAGTGTCTCCTCCTCCAAATGCAAGTGTCTCCTCCTCCAAATGCAAGTGTCTCCTCCTCCAAATGCAAGTGTCTCCTCCTCCAAATGCAAGTGTCTCCTCCTCCAAATGCAAGTGTCTCCTCCTCCAAATGCAAGTGTCTCCTCCTCCAAATGCAAGTGTCTCCTCCAAATGCAAGTGTCTCCTCCAAATGCAAGTGTCTCCTCCAAATGCAAGTGTCTCCTCCAAATGCAAGTGTCTCCTCCAAATGCAAAGTACTCCAAAATGCAAATGCAAATGCAAATGCAAATGCAAATGCAAATGCAAATGCAAATGCAAATGCAAATGCAAATGCAAATGCAAATGCAAATGCAAATGCAAAGTACTCCAAATGCAAATGCAAAGTACTCCAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAAATGCAAAATGCAAAATGCAAAATGCAAAATGCAAAAAAAA	AAAAGTCTGAAACAACATCAAGCGCGTACCCTGCGAAACATCAAGCGCGTACCCTGCGAAACATCAAGCGCGTACCCTGCGAAACATCAAGCGCGTACCCTGCGAAACATCAAGCGCGTACCCTGCGAAACACATCAAGCGCGTACCCTGCGAAACCGACATCAAGCGCGTACCCTGCGAAACCGACATCAAGCGCGTACCCTGCGAAAGAACCGACATCAAGCGCGTACCCTGCGAAACCGACATCAAGCGCGTACCCTGCGAAA	GACACCGCCTTCACTGTCGGAGAACTCAAGAAAGCAATC GACACCGCCCTTCACTGTCGGAGAACTCAAGAAAGCAATC GACACCGCCTTCACTGTCGGAGAACTCAAGAAAGCAATC GACACCGCCTTCACTGTCGGAGAACTCAAGAAAGCAATC GACACCGCCTTCACTGTCGGAGAACTCAAGAAAGCAATC	121 CCACCGCACTGTTTCAAACGCTCGATCCCTCGCTTTTCT 121 CCACCGCACTGTTTCAAACGCTCGATCCCTCGCTTTTCT 121 CCACCGCACTGTTTCAAACGCTCGATCCCTCGCTCTTTCT 122 CCACCGCACTGTTTCAAACGCTCGATCCCTCGCTTTTCT 121 CCACCGCACTGTTTCAAACGCTCGATCCCTCGCTCTTTCT 121 CCACCGCACTGTTTCAAACGCTCGATCCCTCGCTCTTTCT 121 CCACCGCACTGTTTCAAACGCTCGATCCCTCGCTCTTTCT
	4 4 1 1 4 4 1 1 1 1 1 1 1 1 1 1 1 1 1 1	. 81 81 81 81 81	AAAAA

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	CTACCTCATCTGGGACATCATAGCCTCCTGCTTCTA FAd2-D WE CTACCTCATCTGGGACATCATAGCCTCCTGCTTCTA FAd2-D WE CTACCTCATCTGGGACATCATCATAGCCTCCTGCTTCTA FAd2-D (GA316) IMC 129 CTACCTCATCTGGGACATCATCATAGCCTCCTGCTTCTA FAd2-F WE CTACCTCATCTGGGACATCATCATAGCCTCCTGCTTCTA FAd2-F (TA515) Q508 CTACCTCATCTGGGACATCATCATAGCCTCCTGCTTCTA FAd2-F (GA908) Q4275
0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	210 220 230 240 CGTCGCACTTACTTCCCTCTCCTCACCT Fad2-D WE CGTCGCCACCACTTACTTCCTTCCTCCTCCTTCACCT Fad2-D (GA316) IMC 129 CGTCGCCACCACTTACTTCCTTCCTCCTCCTCACCT Fad2-F WE CGTCGCCACCACTTACTTCCTTCCTCCTCCTCACCT Fad2-F (TA515) Q508
0000	250 260 270 280 TCCTACTTCGCCTGGCCTCTCTACTGGGCCTGCCAGG Fad2-D wt TCCTACTTCGCCTGGCTTGCCAGG Fad2-D wt TCCTACTTCGCCTGGCTCTCTACTGGGCCTGCCAGG Fad2-F wt TCCTACTTCGCTGGCTCTTACTGGGCCTGCCAAG Fad2-F (TAS15) Q508 TCCTACTTCGCCTGGCTCTCTACTGGGCCTGCCAAG Fad2-F (TAS15) Q508
000000000000000000000000000000000000000	290 320 320 GCGTCCTAACCGGCGTCTGGGTCATAGCCCACGAGTGFad2-D wt GCGTCCTAACCGGCGTCTGGGTCATAGCCCACAGTGFad2-D (GA316) IMC 129 GCGTCCTAACCGGCGTCTGGGTCATAGCCCACAGTGFad2-F wt GCGTCCTAACCGGCGTCTGGTCATAGCCCACGAGTGFad2-F (TA515) 0508 GCGTCCTAACCGGCGTCTGGGTCATAGCCCACGAGTGFad2-F (TA515) 0508

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	330	340	350 360	١.9.	
321	GGCCACCACGCCTTCA	CGACTAC	G T G G C T G G A C G A G T G G C T G G A C G A	Fad2-D Fad2-D	IMC 129
321	C G G C C A C C A C G C C T T C A G		A G T G G C T T G A C G A C A C A C A C A C A C A C A C	: Fad2-F wt : Fad2-F (TA515)	9050
321	GGCCACCACG	CGACTACC	AGTGGCT	: Fad2-F (GA908)	04275
	370	380	390 400	L 0	
361	CCGTCGGCCTC	CACTCCTT	1	Fad2-D wt (GA316)	IMC 129
361 361		CACTC	CICCICGICCI	Fad2-F	
361	A C C G T C G G T C T C A T C T T C C C T C A T C T T T C C C T C T	CACTCCTT		Fad2-F (TA515)	Q508 Q4275
361		,			
	410	420	430 4.	440	
401	CTTCTCCTGGAAGTAC		CGCCACCATTCCA	A Fad2-D wt	TMC 129
401		GTCATCG	CGCCACCATTCCA	Fad2-F	
401	GTAC	GTCAT	CGCCACCATTCCA	A Fad2-F (TA515)	Q508 04275
401	CTTCTCCTGG	GTCATCG	всемеситеся	radz-r	6.250
	450	460	470 48	480	
441	A C T G G C T C C C	AGACGAAG	TGTTTGTCCCCAA TGTTTGTCCCCAA	G Fad2-D wt G Fad2-D (GA316)	IMC 129
441	A C T G G C T C C C T C G S G A	AGACGAA	TGTTTGTCCCCAA	G Fad2-F wt G Fad2-F (TAS15)	0508
441	C T C C C T	AGACGAA	GTTTGTCCCAA	Fad2 - F	

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PLANTS, SEEDS AND OILS HAVING AN ELEVATED TOTAL
MONOUNSATURATED FATTY ACID CONTENT

IMC 129	IMC 129	IMC 129	IMC 129
Q508	Q508	Q508	Q508
Q4275	Q4275	Q4275	Q4275
Fad2-D wt	Fad2-D wt	Fad2-D wt	Fad2-D wt
Fad2-D (GA316)	Fad2-D (GA316)	Fad2-D (GA316)	Fad2-D (GA316)
Fad2-F wt	Fad2-F wt	Fad2-F wt	Fad2-F wt
Fad2-F (TA515)	Fad2-F (TA515)	Fad2-F (TA515)	Fad2-F (TA515)
Fad2-F (GA908)	Fad2-F (GA908)	Fad2-F (GA908)	Fad2-F (GA908)
AAGAAGTCAGACATCAAGTGGTACGGCAAGTACCTCAACAAAGAGAAGTCAAGTGGTACGGCAAGTACCTCAACAAAGAGAAGTCAAGTACGGCAAGTACCTCAACAAAGAAGAAGTCAACAAGAAGAAGAAGAAGAAGAAGAAGAAGAAGAAGAAG	ACCCTTTGGGACGCACCGTGATGTTAACGGTTCAGTTCA	TCTCGGCTGGCCTTTGTACTTAGCCTTCAACGTCTCGGGGGTCTCGGGGGGTCTCGGGGGGTCTCGGGGGG	AGACCTTACGACGGCGCTTGCTTGCATTTCCACCCAAAGACCTTACGACGCTTGGCTTGCCATTTCCACCCAAAGACGTTACGATTTCCACCCAAAGACGTTACGATTTCCACCCAAAGACGTTACGACGTTGCCATTTCCACCCAAAGACGTTACGACGTTGCCATTTCCACCCAAAGACGTTACGACGCCAAAAGACCTTACGACGCCCAAAAGACCTTACGACGCCCAAAAAGACCTTACGACGCCCAAAAAAAGACCTTACGACGCCCAAAAAAGACCTTACGACGCCCAAAAAAAA
481 481 481 481	521 521 521 521 521 521	561 561 561 561 561	601 601 601 601 601

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PLANTS, SEEDS AND OILS HAVING AN ELEVATED TOTAL
MONOUNSATURATED FATTY ACID CONTENT

IMC 129 Q508 Q4275	IMC 129 Q508 Q4275	IMC 129 Q508 Q4275	IMC 129 Q508 Q4275
Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (TA515) Fad2-F (GA908)	Fad2-D wt. Fad2-D (GA316) Fad2-F wt Fad2-F (TA515)	Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (GA908)	Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (GA908)
650 660 670 680 CGCTCCCATCTACAACGACCGTGAGCGTCTCCAGATATA FE CGCTCCCATCTACAACGACCGTGAGCGTCTCCAGATATA FE CGCTCCCATCTACAACGACCGCGAGCGTCTCCAGATATA FE CGCTCCCATCTACAACGACCGCGAGCGTCTCCAGATATA FE	690 700 710 720 720 720 720 720 720 720 720 720 72	ACCGCTACGCTGTCCAAGGAGTTGCCTCGATGGTCT FacCGCTCGATGGTCT FacCGCTACGCTCGATGGTCT FacCGCTCGATGGTCT FacCGCTCGATGGTCT FacCGCTCGATGGTCT FacCGCTCGCTCGATGGTCT FacCGCTCGCTCGATGGTCT FacCGCTCGCTCGCTCGATGGTCT FacCGCTCGCTCGCTCGATGGTCT Fa	CTTCTACGGAGTTCCTCTTCTGATTGTCAACGGGTTCTT FACTTCTT FACTTTTCTT FACTTCTT FACTTTTCTT FACTTTTCTT FACTTTTTCTT FACTTTTTTTTTT
641 A 641 A 641 A 641 A	681 681 681 681 681	721 T 721 T 721 T 721 T 721 T	761 G 761 G 761 G 761 G

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		•	
IMC 129 Q508 Q4275	IMC 129 Q508 Q4275	IMC 129 Q508 Q4275	IMC 129 Q508 Q4275
) wt) (GA316) ; wt ; (TA515)	wt (GA316) wt (TA515) (GA908)	wt (GA316) wt (TA515) (GA908)	wt (GA316) wt (TA515) (GA908)
Fad2-D Fad2-D Fad2-F Fad2-F Fad2-F	Fad2-D Fad2-D Fad2-F Fad2-F	Fad2-D Fad2-D Fad2-F Fad2-F	Fad2-D Fad2-D Fad2-F Fad2-F Fad2-F
810 820 830 840 AGTTTTGATCACTTGCAGCACACGCATCCTTCCCTG CGTGTTGATCACTTACTTGCAGCACACGCATCCTTCCCTG CGTGTTGATCACTTACTTGCAGCACACGCATCCTTCCCTG CGTGTTGATCACTTACTTGCAGCACACGCATCCTTCCCTG CGTGTTGATCACTTACTTGCAGCACACGCATCCTTCCCTG	CCTCACTATGACTCGTCTGAGTGGGATTGGTTGAGGGGAGCCTCACTATGACTCGTCTGAGTGGGATTGGTTGAGGGGAGGCCTCACTCGTCTGAGTGGGATTGGTTGAGGGGAGGCCTCACTCGTCCGAGTGGGATTGGTTGAGGGGAGGCCTCACTACGATTCGTTCG	CTTTGGCCACCGTTGACAGAGACTACGGAATCTTGAACAA CTTTGGCCACCGTTGACAGAGACTACGGAATCTTGAACAA CTTTGGCTACCGTTGACAGAGACTACGGAATCTTGAACAA CTTTGGCTACCGTTGACAGAGACTACGGAATCTTGAACAA CTTTGGCTACCGTTGACAGACTACGGAATCTTGAACAA CTTTGGCTACCGTTGACAGACTACGGAATCTTGAACAA	GGTCTTCCACATATCACGGACACGCACGTGGCGCATCACGGGTCTTCCACACATCACGGACACGCACG
801 801 801 801	841 841 841 841	881 881 881 881	921 921 921 921

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MONOUNSATURATED FATTY ACID CONTENT

. 6	129	129 1
IMC 129 0508 04275	IMC Q508 Q427	IMC 1 Q508 Q4275
Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (TA515) Fad2-F (GA908)	Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (TA515) Fad2-F (GA908)	Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (TA515)
	Pad2-D wt Fad2-D (G Fad2-F wt Fad2-F (T Fad2-F (G	
TGTTCTCGACCATGCCGCATTATCATGCGATGGAAGCTA TGTTCTCGACCATGCGCATTATCATGCGATGGAAGCTA TGTTCTCCACGATGCCGCATTATCACGCGATGGAAGCTA TGTTCTCCACGATGCCGCATTATCACGCGATGGAAGCTA TGTTCTCCACGATGCCGCATTATCACGCGATGGAAGCTA	1010 1040 1050 1050 1050 1050 1050 GAAGGCGATACTGGGAGAGTATTATCAGTT GAAGGCGATAAACCGATACTGGGAGAGTATTATCAGTT CAAGGCGATAAACCGATACTGGGAGAGTATTATCAGTT CAAGGCGATAAAGCCGATACTGGGAGAGTATTATCAGTT CAAGGCGATAAAGCCGATACTGGGAGAGTATTATCAGTT CAAGGCGATAAAGCCGATACTGGGAGAGTATTATCAGTT	GATGGACGCGGTGGTTAAGGCGATGTGGAGGGAGGCGGATGTGGAGGGGGGGG
961 C 961 C 961 C 961 C	1001 C 1001 C 1001 C 1001 C	1041 C 1041 C 1041 C 1041 C

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	IMC 129 Q508 Q4275	IMC 129 . Q508 Q4275
	116) IM 115) Q5 108) Q4	
	Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (TA515)	Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (TA515) Fad2-F (GA908)
. 0	Fad2-D wt Fad2-D (G) Fad2-F wt Fad2-F (T) Fad2-F (T)	Fad2-D wt Fad2-D (G) Fad2-F wt Fad2-F (T)
1120	CTATGTGGAACCGGACAGGCAAGGTGAGACTATGTGGAACCGGACAGGCAAGGTGAGACTATGTGGAACCGGACAGGCAAGGTGAGACTATGTGGAACCGGACAGGCAAGGTGAGACTATGTGGAACCGGACAGGCAAGGTGAGA	
	1 A G G T A G G G T A G G G G	1 1 1 0 A A A C A A A C A A A A C A A A A A A
1110		1150 A G T T P A G T T P A G T T P A G T T P
		7 A A A A A A A A A A A A A A A A A A A
1100	A A C C C C C C C C C C C C C C C C C C	1140 1150 TTCTGGTACAATAAGTTATGA TTCTGGTACAACAATAAGTTATGA TTCTGGTACAACAATAAGTTATGA TTCTGGTACAACAATAAGTTATGA
1	6 6 7 6 6 6 7 6 6 6 7 6 6 6 6 7 6 6 6 6	G G T A G G G T A A G G G G
		1111 11101 11011
1090	T A T T T A T T A T T A T T A T T A T	1130 1130 1131 1131 131 131 131 131
	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
	A A G G G A A A G G G G G G G G G G G G	A B B B B B B B B B B B B B B B B B B B
•	1081 1081 1081 1081	1121 1121 1121 1121

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1MC129 Q508 Q4275	IMC129 Q508 Q4275	IMC129 0508 04275	1MC129 Q508 Q4275
Fad2-D wt Fad2-D (GA316) 1MC129 Fad2-F wt Fad2-F (TA515) Q508 Fad2-F (GA908) Q4275	Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (TA515) Fad2-F (GA908)	Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (TA515) Fad2-F (GA908)	Fad2-D wt Fad2-P (GA316) Fad2-F wt Fad2-F (TA515) Fad2-F (GA908)
Fad2-D Fad2-D Fad2-F Fad2-F	Fad2-D wt Fad2-D (G Fad2-F wt Fad2-F (T) Fad2-F (G)	Fad2-D wt Fad2-D (G Fad2-F wt Fad2-F (T Fad2-F (G	Fad2-D wt Fad2-D (G) Fad2-F wt Fad2-F (T) Fad2-F (G)
Met Gly Ala Gly Gly Arg Met Gln Val Ser Pro Pro Ser Lya Lya Ser Glu Thr Asp Asn Met Gly Ala Gly Gly Arg Met Gln Val Ser Pro Pro Ser Lya Lya Ser Glu Thr Asp Asn Met Gly Ala Gly Gly Arg Met Gln Val Ser Pro Pro Ser Lya Lya Ser Glu Thr Asp Thr Met Gly Ala Gly Gly Arg Met Gln Val Ser Pro Pro Ser Lya Lya Ser Glu Thr Asp Thr Met Gly Ala Gly Gly Arg Met Gln Val Ser Pro Pro Ser Lya Lya Ser Glu Thr Asp Thr Met Gly Ala Gly Gly Arg Met Gln Val Ser Pro Pro Ser Lya Lya Ser Glu Thr Asp Thr	10 11e Lys Arg Val Pro Cys Glu Thr Pro Pro Phe Thr Val Gly Glu Leu Lys Lys Ala Ile 21 11e Lys Arg Val Pro Cys Glu Thr Pro Pro Phe Thr Val Gly Glu Leu Lys Lys Ala Ile 21 11e Lys Arg Val Pro Cys Glu Thr Pro Pro Phe Thr Val Gly Glu Leu Lys Lys Ala Ile 21 11e Lys Arg Val Pro Cys Glu Thr Pro Pro Phe Thr Val Gly Glu Leu Lys Lys Ala Ile 21 11e Lys Arg Val Pro Cys Glu Thr Pro Pro Phe Thr Val Gly Glu Leu Lys Lys Ala Ile 21 11e Lys Arg Val Pro Cys Glu Thr Pro Pro Phe Thr Val Gly Glu Leu Lys Lys Ala Ile 21 11e Lys Arg Val Pro Cys Glu Thr Pro Pro Phe Thr Val Gly Glu Leu Lys Lys Ala Ile	pro Pro His Cys Phe Lys Arg Ser Ile Pro Arg Ser Phe Ser Tyr Leu Ile Trp Asp Ile 11 Pro Pro His Cys Phe Lys Arg Ser Ile Pro Arg Ser Phe Ser Tyr Leu Ile Trp Asp Ile 11 Pro Pro His Cys Phe Lys Arg Ser Ile Pro Arg Ser Phe Ser Tyr Leu Ile Trp Asp Ile 11 Pro Pro His Cys Phe Lys Arg Ser Ile Pro Arg Ser Phe Ser Tyr Leu Ile Trp Asp Ile 11 Pro Pro His Cys Phe Lys Arg Ser Ile Pro Arg Ser Phe Ser Tyr Leu Ile Trp Asp Ile 11 Pro Pro His Cys Phe Lys Arg Ser Ile Pro Arg Ser Phe Ser Tyr Leu Ile Trp Asp Ile	10 Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro 61 Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro 61 Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro 61 Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro 61 Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro 61 Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro 61 Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro 61 Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro 61 Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro 61 Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro 61 Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro 61 Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro 61 Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro 62 Ile Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro 64 Ile Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Leu Leu Pro His Pro 64 Ile Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Phe Pro Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Thr Tyr Tyr Tyr Val Ala Thr Tyr Tyr Tyr Val Ala Thr Tyr Tyr Tyr Tyr Tyr Tyr Tyr Tyr Tyr Ty

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IMC129 Q508 Q4275	IMC129 Q508 Q4275	IMC129 Q508 Q4275	IMC129 Q508 Q4275
wt (GA316) wt (TA515) (GA908)	Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (GA908)	wt (GA316) wt (TA515) (GA908)	Fad2-D wt Fad2-D (GA316) Fad2-F wt: Fad2-F (TAS15) Fad2-F (GA908)
Fad2-D wt Fad2-D (G Fad2-F wt Fad2-F (T Fad2-F (G		Fad2-D wt Fad2-D (G Fad2-F wt Fad2-F (T)	Fad2-D wt Fad2-D (G/ Fad2-F wt' Fad2-F (T/ Fad2-F (G/
100 81 Leu Ser Tyr Phe Ala Trp Pro Leu Tyr Trp Ala Cys Gln Gly Cys Val Leu Thr Gly Val 81 Leu Ser Tyr Phe Ala Trp Pro Leu Tyr Trp Ala Cys Gln Gly Cys Val Leu Thr Gly Val 81 Leu Ser Tyr Phe Ala Trp Pro Leu Tyr Trp Ala Cys Gln Gly Cys Val Leu Thr Gly Val 81 Leu Ser Tyr Phe Ala Trp Pro Leu Tyr Trp Ala Cys Gln Gly Cys Val Leu Thr Gly Val 81 Leu Ser Tyr Phe Ala Trp Pro Leu Tyr Trp Ala Cys Gln Gly Cys Val Leu Thr Gly Val 81 Leu Ser Tyr Phe Ala Trp Pro Leu Tyr Trp Ala Cys Gln Gly Cys Val Leu Thr Gly Val	110 120 170 170 170 170 170 170 170 170 170 17	130 121 Thr Val Gly Leu Ile Phe His Ser Phe Leu Leu Val Pro Tyr Phe Ser Trp Lys Tyr Ser 121 Thr Val Gly Leu Ile Phe His Ser Phe Leu Leu Val Pro Tyr Phe Ser Trp Lys Tyr Ser 121 Thr Val Gly Leu Ile Phe His Ser Phe Leu Leu Val Pro Tyr Phe Ser Trp Lys Tyr Ser 121 Thr Val Gly Leu Ile Phe His Ser Phe Leu Leu Val Pro Tyr Phe Ser Trp Lys Tyr Ser 121 Thr Val Gly Leu Ile Phe His Ser Phe Leu Leu Val Pro Tyr Phe Ser Trp Lys Tyr Ser	150 141 His Arg Arg His His Ser Asn Thr Gly Ser Leu Glu Arg Asp Glu Val Phe Val Pro Lys 141 His Arg Arg His His Ser Asn Thr Gly Ser Leu Glu Arg Asp Glu Val Phe Val Pro Lys 141 His Arg Arg His His Ser Asn Thr Gly Ser Leu Glu Arg Asp Glu Val Phe Val Pro Lys 141 His Arg Arg His His Ser Asn Thr Gly Ser Leu Glu Arg Asp Glu Val Phe Val Pro Lys 141 His Arg Arg His His Ser Asn Thr Gly Ser Leu Glu Arg Asp Glu Val Phe Val Pro Lys
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PLANTS, SEEDS AND OILS HAVING AN ELEVATED TOTAL
MONOUNSATURATED FATTY ACID CONTENT

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IMC129 Q508 Q4275	IMC129 Q508 Q4275	IMC129 Q508 Q4275	IMC129 Q508 Q4275
Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (TA515) Fad2-F (GA908)	Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (TA515)	Fad2-D wt. Fad2-D (GA316) Fad2-F wt. Fad2-F (TA515)	Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (TA515)
Lys Lys Ser Asp Ile Lys Trp Tyr Gly Lys Tyr Leu Asn Asn Pro Leu Gly Arg Thr Val Lys Lys Ser Asp Ile Lys Trp Tyr Gly Lys Tyr Leu Asn Asn Pro Leu Gly Arg Thr Val Lys Lys Ser Asp Ile Lys Trp Tyr Gly Lys Tyr Leu Asn Asn Pro Leu Gly Arg Thr Val Lys Lys Ser Asp Ile Lys Trp Tyr Gly Lys Tyr His Asn Asn Pro Leu Gly Arg Thr Val Lys Lys Ser Asp Ile Lys Trp Tyr Gly Lys Tyr Leu Asn Asn Pro Leu Gly Arg Thr Val	Met Leu Thr Val Gln Phe Thr Leu Gly Trp Pro Leu Tyr Leu Ala Phe Asn Val Ser Gly Met Leu Thr Val Gln Phe Thr Leu Gly Trp Pro Leu Tyr Leu Ala Phe Asn Val Ser Gly Met Leu Thr Val Gln Phe Thr Leu Gly Trp Pro Leu Tyr Leu Ala Phe Asn Val Ser Gly Met Leu Thr Val Gln Phe Thr Leu Gly Trp Pro Leu Tyr Leu Ala Phe Asn Val Ser Gly Met Leu Thr Val Gln Phe Thr Leu Gly Trp Pro Leu Tyr Leu Ala Phe Asn Val Ser Gly Met Leu Thr Val Gln Phe Thr Leu Gly Trp Pro Leu Tyr Leu Ala Phe Asn Val Ser Gly	Arg Pro Tyr Asp Gly Gly Phe Ala Cys His Phe His Pro Asn Ala Pro Ile Tyr Asn Asp Arg Pro Tyr Asp Gly Gly Phe Ala Cys His Phe His Pro Asn Ala Pro Ile Tyr Asn Asp Arg Pro Tyr Asp Gly Gly Phe Ala Cys His Phe His Pro Asn Ala Pro Ile Tyr Asn Asp Arg Pro Tyr Asp Gly Gly Phe Ala Cys His Phe His Pro Asn Ala Pro Ile Tyr Asn Asp Arg Pro Tyr Asp Gly Gly Phe Ala Cys His Phe His Pro Asn Ala Pro Ile Tyr Asn Asp Arg Pro Tyr Asp Gly Gly Phe Ala Cys His Phe His Pro Asn Ala Pro Ile Tyr Asn Asp	Arg Glu Arg Leu Gln Ile Tyr Ile Ser Asp Ala Gly Ile Leu Ala Val Cys Tyr Gly Leu Arg Glu Arg Leu Gln Ile Tyr Ile Ser Asp Ala Gly Ile Leu Ala Val Cys Tyr Gly Leu Arg Glu Arg Leu Gln Ile Tyr Ile Ser Asp Ala Gly Ile Leu Ala Val Cys Tyr Gly Leu Arg Glu Arg Leu Gln Ile Tyr Ile Ser Asp Ala Gly Ile Leu Ala Val Cys Tyr Gly Leu Arg Glu Arg Leu Gln Ile Tyr Ile Ser Asp Ala Gly Ile Leu Ala Val Cys Tyr Gly Leu Arg Glu Arg Leu Gln Ile Tyr Ile Ser Asp Ala Gly Ile Leu Ala Val Cys Tyr Gly Leu
161 161 161 161 161	18 18 18 18 18	.201 201 201 201 201	221 221 221 221 221

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IMC129 Q508 Q4275	IMC129 Q508 Q4275	1MC129 Q508 Q4275	1MC129 Q508 Q4275
Fad2-D wt Fad2-D (GA316) IMC129 Fad2-F wt Fad2-F (TA515) Q508 Fad2-F (GA908) Q4275	wt (GA316) wt (TAS15) (GA908)	A316) A515) A908)	Fad2-D wt. Fad2-F wc. Fad2-F wc. Fad2-F (TA515) Q508 Fad2-F (GA908) Q4275
Fad2-D wt Fad2-D (G Fad2-F wt Fad2-F (T	Fad2-D wt Fad2-D (G Fad2-F wt Fad2-F (T Fad2-F (G	Fad2-D wt Fad2-D (G Fad2-F wt Fad2-F (T Fad2-F (G	Fad2-D wt. Fad2-D (G Fad2-F wt. Fad2-F (T
Tyr Arg Tyr Ala Ala Val Gln Gly Val Ala Ser Met Val Cys Phe Tyr Gly Val Pro Leu Fry Arg Tyr Ala Ala Ala Gln Gly Val Ala Ser Met Val Cys Phe Tyr Gly Val Pro Leu Fre Arg Tyr Ala Ala Ala Gln Gly Val Ala Ser Met Val Cys Phe Tyr Gly Val Pro Leu Fre Arg Tyr Ala Ala Gln Gly Val Ala Ser Met Val Cys Phe Tyr Gly Val Pro Leu Fre Arg Tyr Ala Ala Gln Gly Val Ala Ser Met Val Cys Phe Tyr Gly Val Pro Leu Fre Arg Tyr Ala Ala Gln Gly Val Ala Ser Met Val Cys Phe Tyr Gly Val Pro Leu Fre Arg Tyr Ala Ala Gln Gly Val Ala Ser Met Val Cys Phe Tyr Gly Val Pro Leu Fre	Leu Ile Val Asn Gly Phe Leu Val Leu Ile Thr Tyr Leu Gln His Thr His Pro Ser Leu Leu Ile Val Asn Gly Phe Leu Val Leu Ile Thr Tyr Leu Gln His Thr His Pro Ser Leu Eu Ile Val Asn Gly Phe Leu Val Leu Ile Thr Tyr Leu Gln His Thr His Pro Ser Leu Eu Ile Val Asn Gly Phe Leu Val Leu Ile Thr Tyr Leu Gln His Thr His Pro Ser Leu Eue Ile Val Asn Gly Phe Leu Val Leu Ile Thr Tyr Leu Gln His Thr His Pro Ser Leu Eue Ile Val Asn Gly Phe Leu Val Leu Ile Thr Tyr Leu Gln His Thr His Pro Ser Leu Fueu Ile Val Asn Gly Phe Leu Val Leu Ile Thr Tyr Leu Gln His Thr His Pro Ser Leu Fueu Ile	Pro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Fro His Tyr Asp Ser Ser Glu Trp Asp Trp Trp Trp Trp Trp Trp Trp Trp Trp Tr	Asp Tyr Gly Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Basp Tyr Gly Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Basp Tyr Gly Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Basp Tyr Gly Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Basp Tyr Gly Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Basp Tyr Glu Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Hasp Tyr Glu Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His His Fash Tyr Glu Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Fash Tyr Glu Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Fash Tyr Glu Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Fash Tyr Glu Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Fash Tyr Glu Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Fash Tyr Glu Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Fash Tyr Glu Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Fash Tyr Glu Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Fash Tyr Glu Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Fash Tyr Glu Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His His Fash Tyr Glu Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Fash Tyr Glu Ile Leu Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His
241 241 241 241	261 261 261 261 261	281 281	301 301 301 301 301

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Applicant(s): Dharma R. Kodali et al.
PLANTS, SEEDS AND OILS HAVING AN ELEVATED TOTAL
MONOUNSATURATED FATTY ACID CONTENT

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IMC129 Q508 Q4275	IMC129 Q508 Q4275	IMC129 Q508 Q4275	IMC129 QSOB Q4275
Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (TA515) Fad2-F (GA908)	Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (TA515) Fad2-F (GA908)	Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (TA515) Fad2-F (GA908)	Fad2-D wt Fad2-D (GA316) Fad2-F wt Fad2-F (GA908)
Fad2-D Fad2-D Fad2-F Fad2-F Fad2-F	Fad2-D wt Fad2-D (G Fad2-F wt Fad2-F (T	Fad2-D wt Fad2-D (G Fad2-F wt Fad2-F (T	Fad2-D Fad2-D Fad2-F Fad2-F
His Ala Met Glu Ala Thr Lys Ala Ile Lys Pro Ile His Ala Met Glu Ala Thr Lys Ala Ile Lys Pro Ile His Ala Met Glu Ala Thr Lys Ala Ile Lys Pro Ile His Ala Met Glu Ala Thr Lys Ala Ile Lys Pro Ile His Ala Met Glu Ala Thr Lys Ala Ile Lys Pro Ile His Ala Met Glu Ala Thr Lys Ala Ile Lys Pro Ile	Gly Thr Pro Val Val Lys Ala Met Trp Arg Glu Ala Gly Thr Pro Val Val Lys Ala Met Trp Arg Glu Ala Gly Thr Pro Val Val Lys Ala Met Trp Arg Glu Ala Gly Thr Pro Val Val Lys Ala Met Trp Arg Glu Ala Gly Thr Pro Val Val Lys Ala Met Trp Arg Glu Ala Gly Thr Pro Val Val Lys Ala Met Trp Arg Glu Ala Gly Thr Pro Val Val Lys Ala Met Trp Arg Glu Ala	Asp Arg Gln Gly Glu Lys Lys Gly Val Phe Trp Tyr Asp Arg Gln Gly Glu Lys Lys Gly Val Phe Trp Tyr Asp Arg Gln Gly Glu Lys Lys Gly Val Phe Trp Tyr Asp Arg Gln Gly Glu Lys Lys Gly Val Phe Trp Tyr Asp Arg Gln Gly Glu Lys Lys Gly Val Phe Trp Tyr Asp Arg Gln Gly Glu Lys Lys Gly Val Phe Trp Tyr Asp Arg Gln Gly Glu Lys Lys Gly Val Phe Trp Tyr	
Leu Phe Ser Thr Met Pro His Tyr Leu Phe Ser Thr Met Pro His Tyr Isl Leu Phe Ser Thr Met Pro His Tyr Isl Leu Phe Ser Thr Met Pro His Tyr Isl Leu Phe Ser Thr Met Pro His Tyr Isl Leu Phe Ser Thr Met Pro His Tyr	Leu Gly Glu Tyr Tyr Gln Phe Asp 141 Leu Gly Glu Tyr Tyr Gln Phe Asp	61 Lys Glu Cys Ile Tyr Val Glu Pro 61 Lys Glu Cys Ile Tyr Val Glu Pro	81 Asn Asn Lys Leu ter 81 Asn Asn Lys Leu ter 81 Asn Asn Lys Leu ter 181 Asn Asn Lys Leu ter 181 Asn Asn Lys Leu ter

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PLANTS, SEEDS AND OILS HAVING AN ELEVATED TOTAL

MONOUNSATURATED FATTY ACID CONTENT

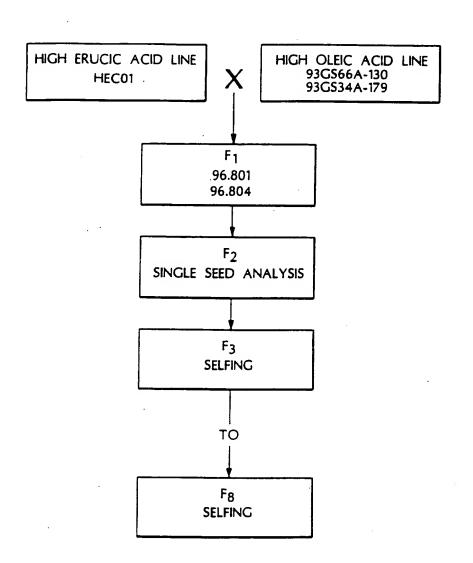


FIG. 4